

09/195,368

**Amendments to the specification:**

Please enter into the specification the enclosed substitute Sequence Listing which includes 9 pages and SEQ ID NO:s 1-15.

On page 8, lines 30-33, in the Brief Description of the Drawings, please amend the text as follows:

Figure 2 shows an alignment and comparison of extracellular amino acid sequence of DNA19355 polypeptide (SEQ ID NO:9) with human Apo-2L (SEQ ID NO:11), Fas/Apo1 ligand (CD95L) (SEQ ID NO:12), TNF-alpha (SEQ ID NO:10) and LT- $\alpha$  (SEQ ID NO:13); the respective amino acid identities (%) are approximately 19.8, 19.0, 20.6, and 17.5.

On page 9, lines 22-24, in the Brief Description of the Drawings, please amend the text as follows:

Figure 7 shows an alignment and comparison of the amino acid sequences for human GITR (hGITR) (SEQ ID NO:14) and murine GITR (mGITR) (SEQ ID NO:15). The three cysteine rich domains (CRD1, CRD2, and CRD3) and transmembrane region (TM) are shown.